RAW SEQUENCE LISTING

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Application Serial Number: 10/549,782
Source: PCT
Date Processed by STIC: 10/03/2005

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RAW SEQUENCE LISTING DATE: 10/03/2005
PATENT APPLICATION: **US/10/549,782** TIME: 14:29:44

Input Set : **A:\Sequence listing - 09600-00035-US.txt**
Output Set: **N:\CRF4\10032005\J549782.raw**

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      3 <110> APPLICANT: Egorova, Ksenia
      4       Antranikian, Garabed
      5       Trauthwein, Harald
      6       Verseck, Stefan
      7       Dingerdissen, Uwe
      9 <120> TITLE OF INVENTION: THERMALLY STABLE AMIDASES
     11 <130> FILE REFERENCE: 09600-00035-US
C--> 13 <140> CURRENT APPLICATION NUMBER: US/10/549,782
C--> 13 <141> CURRENT FILING DATE: 2005-09-19
     13 <150> PRIOR APPLICATION NUMBER: PCT/EP2004/001430
     14 <151> PRIOR FILING DATE: 2004-02-16
     16 <150> PRIOR APPLICATION NUMBER: DE 103 12 842.5
     17 <151> PRIOR FILING DATE: 2003-03-21
     19 <160> NUMBER OF SEQ ID NOS: 4
     21 <170> SOFTWARE: PatentIn version 3.3
     23 <210> SEQ ID NO: 1
     24 <211> LENGTH: 9
     25 <212> TYPE: PRT
     26 <213> ORGANISM: Pseudonocardia thermophila
     28 <400> SEQUENCE: 1
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     35 <211> LENGTH: 14
     36 <212> TYPE: PRT
     37 <213> ORGANISM: Pseudonocardia thermophila
     39 <400> SEQUENCE: 2
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     45 <210> SEQ ID NO: 3
     46 <211> LENGTH: 509
     47 <212> TYPE: PRT
     48 <213> ORGANISM: Pseudonocardia thermophila
     50 <400> SEQUENCE: 3
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     60 Leu Gly His Arg Pro Ala Arg Leu Val Glu Gln Gly Arg Arg Ala Val
     61        35                  40                  45
     64 Arg Gly Gly Gly Pro His Pro Ala Ala Ala Leu Val Asp Pro Pro Arg
     65    50                  55                  60
     68 Arg Arg Gly Arg Gln Ala Arg Arg Val Gly Gly Ala Asp Leu Asp His
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69 65                  70                  75                  80
72 Arg Asn Leu Gly Gly Ala Ala Arg Arg Pro His Gly Arg Gly Gln Asp
73                  85                  90                  95
76 Asn Val Ala Val Ala Gly Val Pro Met Met Asn Gly Ser Arg Thr Leu
77             100                 105                 110
80 Glu Gly Phe Val Pro Ala Glu Asp Ala Thr Val Val Ser Arg Leu Leu
81         115                 120                 125
84 Ala Ala Gly Ala Thr Ile Ala Gly Lys Ser Val Cys Glu Asp Leu Cys
85     130                 135                 140
88 Phe Ser Gly Gly Ser His Thr Ser Lys Pro Gly Thr Val His Asn Pro
89 145                 150                 155                 160
92 Trp Asp Met Ser Arg Ser Ala Gly Gly Ser Ser Ser Gly Ser Gly Ala
93                  165                 170                 175
96 Leu Val Ala Ala Gly Glu Val Asp Met Ala Ile Gly Gly Asp Gln Gly
97             180                 185                 190
100 Gly Ser Ile Arg Ile Pro Ser Ala Tyr Cys Gly Thr Val Gly His Lys
101         195                 200                 205
104 Pro Thr His Gly Leu Val Pro Tyr Thr Gly Gly Phe Pro Ile Glu Gln
105     210                 215                 220
108 Ser Ile Asp His Leu Gly Pro Ile Thr Arg Thr Val Ala Asp Ala Ala
109 225                 230                 235                 240
112 Leu Met Leu Ser Val Ile Ala Gly Arg Asp Gly Leu Asp Pro Arg Gln
113                 245                 250                 255
116 Pro Asp Val Val Glu Val Gln Asp Tyr Val Gly Ala Leu Ala Glu Ser
117             260                 265                 270
120 Val Ser Gly Leu Arg Ile Gly Val Leu Gln Glu Gly Phe Gly His Pro
121         275                 280                 285
124 Asn Ser Glu Pro Glu Val Asp Asp Thr Val Arg Ala Ala Val Gly Thr
125     290                 295                 300
128 Leu Arg Glu Ala Gly Phe Thr Val Glu Asp Val Ser Val Pro Trp His
129 305                 310                 315                 320
132 Leu His Ala Thr Ala Ile Trp Asp Val Leu Ala Thr Glu Gly Gly Leu
133                 325                 330                 335
136 Trp Gln Met Val Glu Gly Asn Ala Tyr Gly Met His Trp Lys Gly His
137             340                 345                 350
140 Tyr Asp Pro Ser Leu Ile Ala Tyr Tyr Gly Arg Lys Trp Arg Glu Asp
141         355                 360                 365
144 Pro Ala Gln Phe Ser Glu Thr Val Lys Leu Val Ala Leu Ala Gly Arg
145     370                 375                 380
148 Tyr Ala Leu Thr Thr Gln Tyr Gly Arg His Tyr Ala Met Ala Arg Asn
149 385                 390                 395                 400
152 Leu Ala Pro Lys Leu Val Ala Ala Tyr Asp Ala Ala Leu Ser Asn Tyr
153                 405                 410                 415
156 Asp Val Leu Val Met Pro Thr Leu Pro Met Arg Ala Thr Val Leu Pro
157             420                 425                 430
160 Gly Pro Asp Ala Pro Val Glu Glu Ile Leu Ala Arg Gly Leu Glu Met
161         435                 440                 445
164 Leu Ala Asn Thr Ala Pro Phe Asp Val Thr Gly His Pro Ala Cys Ser
165     450                 455                 460
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168 Val Pro Ala Gly Leu Ala Asp Gly Leu Pro Val Gly Leu Met Ile Val
169 465                 470                 475                 480
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177             500                 505
180 <210> SEQ ID NO: 4
181 <211> LENGTH: 1527
182 <212> TYPE: DNA
183 <213> ORGANISM: Pseudonocardia thermophila
185 <400> SEQUENCE: 4
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190 gtggaacaag gtcgaagagc tgtacgcgga ggtggccccc accccgccgc agcgctcgtg    180
192 gacccgcccc gacgccgagg acgacaagct cggcgcgtgg gcggtgcaga cctcgatcac    240
194 cgaaacctcg gaggggccgc tcgccggccg cacggtcgcg gtcaggacaa cgtcgccgtc    300
196 gccggcgtgc cgatgatgaa cggatcgcgg accctggagg gcttcgtccc cgccgaggac    360
198 gccaccgtcg tgtcccggct gctcgccgcc ggcgcgacca tcgccggcaa gtcggtgtgc    420
200 gaggacctgt gcttctccgg cggctcgcac accagcaagc cgggaaccgt gcacaacccg    480
202 tgggacatga gccgctcggc gggtggttcc tcgagcggta gcggggcgct ggtggccgcc    540
204 ggcgaggtcg acatggccat cggcggtgac cagggcgggt cgatccgcat cccgtcggcg    600
206 tactgcggca ccgtcgggca caagccgacc cacggcctgg tgccgtacac cggaggcttc    660
208 ccgatcgagc agtcgatcga ccacctcggc ccgatcaccc gcaccgtcgc cgacgccgcg    720
210 ctcatgctgt cggtgatcgc cggccgcgac gggctcgacc cgcgccagcc cgacgtcgtc    780
212 gaggtgcagg actacgtcgg cgcgctggcc gagtcggtga gcgggctgcg catcggcgtc    840
214 ctgcaggagg gcttcggtca ccccaactcc gagcccgagg tcgacgacac cgtgcgcgct    900
216 gctgtcggga cactgcgcga ggccggcttc accgtcgagg acgtctcggt cccgtggcac    960
218 ctgcacgcca cggccatctg ggacgtgctc gccaccgagg gcggcctgtg gcagatggtc   1020
220 gagggcaacg cctacggcat gcactggaag ggccactacg acccgtcgct catcgcctac   1080
222 tacgggcgca agtggcggga ggacccggcg cagttctccg agacggtcaa gctcgtcgcc   1140
224 ctcgccggcc gctacgcgct cacgacccag tacgggcgcc actacgcgat ggcgcgcaac   1200
226 ctcgccccga agctggtcgc ggcctacgac gcggcgctgt cgaactacga cgtgctcgtc   1260
228 atgccgacgc tgccgatgcg cgccaccgtg ctgcccggcc cggacgcgcc cgtcgaggag   1320
230 atcctcgcca ggggcctgga gatgctggcc aacaccgcac cgttcgacgt caccgggcac   1380
232 ccggcctgct ctgtcccggc cggcctcgcc gacggtctcc cggtcggcct gatgatcgtc   1440
234 ggcaagcact tcgacgacgc caccgtgctg aaggtggcga acgccttcga acaggccgtc   1500
236 ggcggcttcc cgacccccgc tgcctga                                      1527
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VERIFICATION SUMMARY DATE: 10/03/2005
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L:13 M:271 C: Current Filing Date differs, Replaced Current Filing Date